



Information Disclosure Statement By Applicant (Use Several Sheets if Necessary)	Atty Docket No. ICYTP003X1C2 Applicant: Koleszar et al. Filing Date Nov. 3. 2003	Application No.: 10/700,792 Group Not yet assigned
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Examiner Initial	No.	Patent No.	Date	Patentee	Class	Sub-class	Filing Date
M.D.P.	A	5,970,500	10/1999	Sabatini, et al.			
	B	5,953,727	09/1999	Maslyn, et al.			
	C	6,023,659 A	02/2000	Seilhamer, et al.			
	D	5,418,944	05/23/95	DiPace, et al.			
	E	5,706,498	1/6/98	Fujimiya et al.			
	F	5,523,208	6/7/96	Kohler et al.			
	G	5,840,484	11/24/98	Seilhamer et al.			
	H	5,701,256	12/23/97	Marr et al.			
	I	5,681,285	10/28/97	Ford et al.			
M.D.P.	J	5,840,562	11/24/98	Diep et al.			

Foreign Patent or Published Foreign Patent Application

Examiner Initial	No.	Document No.	Publication Date	Country or Patent Office	Class	Sub-class	Translation	
							Yes	No
M.D.P.	A	EP 0 848 067 A2	06/17/98	European	C12Q 1	68	X	
M.D.P.	B	WO/96/23078	01.08.96	PCT			X	

Other Documents

Examiner Initial	No.	Author, Title, Date, Place (e.g. Journal) of Publication
M.D.P.	A	Duret, et al., "HOVERGEN: a database of homologous vertebrate genes", Nucleic Acids Research, Received February 17, 1994, Revised and Accepted May 10, 1994, Vol. 22, No. 12, pp. 2360-2365.
M.D.P.	B	Larsen, et al., "The ribosomal database project", Nucleic Acids Research, 1993, Vol. 21, no. 13, pp. 3021-3023.
M.D.P.	C	D.W. Smith, et al., "Biocomputing: Informatics Genome Projects", Academic Press, Inc., Department of Biology and Center for Molecular Genetics, University of California, 1994, pp. 51-85.
M.D.P.	D	Parsons, et al., "Clustering CDNA Sequences", CABIOS, 8(5) 461-466.
M.D.P.	E	Henikoff, Steven, "Comparative Sequence Analysis: Finding Genes", Academic Press, Inc., 1994, pp. 867-111.
Examiner	<i>Michael D. Plummer</i> Date Considered 7/20/2006	

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Patent No. 1449 (Modified) Information Disclosure Statement By Applicant (Use Several Sheets if Necessary)	Atty Docket No. ICYTP003X1C2 Applicant: Koleszar et al. Filing Date Nov. 3. 2003	Application No.: 10/700,792 Group Not yet assigned
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Other Documents

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M.D.P.	F	Schena, et al., "Parallel human genome analysis: Microarray-based expression monitoring of 1000 genes", Proc. Natl. Acad. Sci. USA, Vol. 93, October 1996, pp. 10614-10619.
	G	Cuticchia, et al., "CMAP: contig mapping and analysis package, a relational database for chromosome reconstruction", Department of Genetics, The University of Georgia, Vol. 8, No. 5, 1992, pp. 467-474.
	H	Fickett, J.W. "Finding genes by computer: the state of the art" TIG Vol. 12, No.8, pp. 316-320. (9/1996)
	I	Honda, et al., "An Object Model for Genome Information at All Levels of Resolution", January 1993, Proceedings of the Twenty-sixth Hawaii International Conference on System Sciences, Vol. I, pp. 564-573.
	J	Garner, "Engineering ins Genomiocs/ spl minus/ Automating the Gemone Center", May 1994, IEEE Engineering in Medicine and Biology Magazine, Vol. 13, No. 2, pp. 281-283.
	K	PathoSeq. 3.0 December 97 Release Notes
	L	The Institute for Genomic Research, TIGR Database. Internet - http://www.tigr.org (Downloaded December 23, 1997).
	M	National Center for Biotechnology Information (Index). Internet- http://www3.ncbi.nlm.nih.gov/ (Downloaded December 23, 1997).
	N	National Center for Biotechnology Information; ENTREZ. Internet- http://www3.ncbi.nlm.nih.gov/entrez (Downloaded December 23, 1997).
	O	National Center for Biotechnology Information; UniGene. Internet- http://www3.ncbi.nlm.nih.gov/unigene (Downloaded December 23, 1997).
	P	National Center for Biotechnology Information; GenBank. Internet- http://www3.ncbi.nlm.nih.gov/web/genbank (Downloaded December 23, 1997).
	Q	MAGPIE; Automated Genome Project Investigation Environment. Internet- http://www.mcs.anl.gov/home/gaasterl/magpie.html (Downloaded December 23, 1997).
M.D.P.	R	Stanford University, Stanford Genomic Resources. Internet- http://genome-www.stanford.edu , September 1996 and December 20, 1997.
Examiner	<i>Michael D. Klein</i> Date Considered <i>7/20/2006</i>	

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Examiner Initial	No.	Author, Title, Date, Place (e.g. Journal) of Publication
M.D.P.	S	Incyte Pharmaceuticals, LIFESEQ Version 4.2 Release Notes and Physical Data Model, October 1996.
	T	Incyte Pharmaceuticals, LIFESEQ Version 4.1 Release Notes and Physical Data Model, July 1996.
	U	Incyte Pharmaceuticals, LIFESEQ Version 3.4 Release Notes, January 1996.
	V	Incyte Pharmaceuticals, LIFESEQ Version 2.5 Release Notes, June 1995.
	W	Incyte Pharmaceuticals, LIFESEQ Version 3.0 Release Notes, September 1995.
	X	Incyte Pharmaceuticals, LIFESEQ Version 4.0 Release Notes, April 1996.
	Y	Incyte Pharmaceuticals, <u>Introduction to the LIFESEQ Database</u> , Version 3.4, January 1996.
	Z	Incyte Pharmaceuticals, <u>LIFESEQ Training Manual</u> , Version 4.1, July 1996.
	AA	Green, et al., "Ancient Conserved Regions in New Gene Sequences and the Protein Databases", <u>Science</u> , v.259, n.5102, pgs. 1711-1716 (1993)
	AB	Waldrop, et al., "Online Archives Lets Biologist Interrogate the Genome", <u>Science</u> , v. 269, n. 5229, pgs. 1356-1358 (1995)
	AC	No Author, "Incyte Serves Up Information, part I", <u>In Vivo the Business & Medicine Report</u> , p.32, ISSN: 0258-851X (1996).
	AD	Martin, et al., "Accessing Genetics Databases", <u>Database</u> , v.17, n. 1, pg. 51(8) (1994).
	AE	Matsubara, et al., "Identification of New Genes by Systematic Analysis of cDNAs and Database Construction", <u>Current Opinion in Biology</u> , Vol. 4, pp. 672-677 (1993).
	AF	Lew, K., "GDB 6.0 Goals", http://gdb.gbdnet.ad.jp/gdb/docs/gdb6-goals.html , pp. 1-5 (March 3, 1996).
	AG	Kanehisa, M. "Toward Pathway Engineering: A New Database Old Genetic and Molecular Pathways." <u>Science and Technology Japan</u> , 1995, No. 59, pages 34-38.
M.D.P.	AH	Gaasterland, T. and Sensen, C. "Using Multiple Tools for Automated Genome Interpretation in an Integrated Environment", <u>Trends In Genetics</u> , January 1996.
Examiner	<i>Michael D. Phan</i> Date Considered <i>7/20/2006</i>	

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Form 1449 (Modified)

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M.D.P.	AI	http://wehih.wehi.edu.au/ (1995).
	AJ	Keele et al. "A Conceptual Database Model for Genomic Research", Journal Of Computational Biology, Vol. 1, No. 1 pp. 65-76 (1994)
	AK	Adams, M.D. et al. "Complementary DNA Sequencing: Expressed Sequence Tags and Human Genome Project", Science, 21 June 1991, Vol. 252, page 1651-1656.
	AL	Keele et al., "A Conceptual Database Model for Genomic Research", 1994, Journal of Computational Biology, Vol. 1, No. 1, pp. 65-76.
	AM	Mount, D.W. et al., "A Genomic Database for Escherichia coli: Total Information on a Given Organism" Academic Press BIOCOMPUTING 1994, pp. 249-266.
M.D.P.	AN	Yoshida, K. et al., "A Primer on Rapid Prototyping of Genomic Databases in Prolog." Academic Press BIOCOMPUTING, 1994, pp. 233-247.
Examiner	<i>Michael D. Pfen</i>	
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